

SEQUENCE LISTING

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 BRUNNER, Erich
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 KRAMPS, Thomas
 PETER, Oliver

<120> ESSENTIAL DOWNSTREAM COMPONENT OF THE WINGLESS SIGNALING PATHWAY AND THERAPEUTIC AND DIAGNOSTIC APPLICATIONS BASED THEREON

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<130> Q60361
<140> 09/915,543
<141> 2001-07-27
<150> 60/221,502
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- Gly Ser Met Gly Leu Lys Asn Gly Ala Gly Asn Gly Ala Lys Gly Lys 85 90 95
- Gly Lys Arg Glu Arg Ser Ile Ser Ala Asp Ser Phe Asp Gln Arg Asp 100 105 110
- Pro Gly Thr Pro Asn Asp Asp Ser Asp Ile Lys Glu Cys Asn Ser Ala 115 120 125
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- Pro Ser Asn Ala Thr Ala Pro Arg Ser Ser Thr Pro Ser His Gly Gln 145 150 155 160
- Thr Thr Ala Thr Glu Pro Thr Pro Ala Gln Lys Thr Pro Ala Lys Val 165 170 175
- Val Tyr Val Phe Ser Thr Glu Met Ala Asn Lys Ala Ala Glu Ala Val 180 185 190
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- Phe Gly Glu His Pro Gln Gln Glu Tyr Gly Met Gly Pro Arg Pro Phe 770 775 780
- Leu Pro Met Ser Gln Gly Pro Gly Ser Asn Ser Gly Leu Arg Asn Leu 785 790 795 800
- Arg Glu Pro Ile Gly Pro Asp Gln Arg Thr Asn Ser Arg Leu Ser His 805 810 815
- Met Pro Pro Leu Pro Leu Asn Pro Ser Ser Asn Pro Thr Ser Leu Asn 820 825 830
- Thr Ala Pro Pro Val Gln Arg Gly Leu Gly Arg Lys Pro Leu Asp Ile 835 840 845
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- Ser Pro Thr Met His Gln Val Gln Ser Pro Met Leu Gly Ser Pro Ser 865 870 875 880
- Gly Asn Leu Lys Ser Pro Gln Thr Pro Ser Gln Leu Ala Gly Met Leu 885 890 895
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- Ser Ala Ala Ser Pro Val His Leu Lys Ser Pro Ser Leu Pro Ala 915 920 925
- Pro Ser Pro Gly Trp Thr Ser Ser Pro Lys Pro Pro Leu Gln Ser Pro 930 935 940
- Gly Ile Pro Pro Asn His Lys Ala Pro Leu Thr Met Ala Ser Pro Ala 945 950 955 960

- Met Leu Gly Asn Val Glu Ser Gly Gly Pro Pro Pro Pro Thr Ala Ser 965 970 975
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- Tyr Thr Met Pro Pro Glu Pro Thr Leu Ser Gln Asn Pro Leu Ser Ile 995 1000 1005
- Met Met Ser Arg Met Ser Lys Phe Ala Met Pro Ser Ser Thr Pro 1010 1015 1020
- Leu Tyr His Asp Ala Ile Lys Thr Val Ala Ser Ser Asp Asp Asp 1025 1030 1035
- Ser Pro Pro Ala Arg Ser Pro Asn Leu Pro Ser Met Asn Asn Met 1040 1045 1050
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- Gln Pro Leu Ser His Ser Asn Gln Met Pro Ser Pro Asn Ala Val 1085 1090 1095
- Gly Pro Asn Ile Pro Pro His Gly Val Pro Met Gly Pro Gly Leu 1100 1105 1110
- Met Ser His Asn Pro Ile Met Gly His Gly Ser Gln Glu Pro Pro 1115 1120 1125
- Met Val Pro Gln Gly Arg Met Gly Phe Pro Gln Gly Phe Pro Pro 1130 1135 1140
- Val Gln Ser Pro Pro Gln Gln Val Pro Phe Pro His Asn Gly Pro 1145 1150 1155
- Ser Gly Gly Gln Gly Ser Phe Pro Gly Gly Met Gly Phe Pro Gly 1160 1165 1170
- Glu Gly Pro Leu Gly Arg Pro Ser Asn Leu Pro Gln Ser Ser Ala

1175 1180 1185

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Ala Ser Phe Leu Gly Gln Gln Gly Arg Val Ile Trp Lys Pro Leu Ser 35 40 45

Glu Glu Leu Arg Asp Gln Gly Ala Asp Ala Ala Gly Gly Pro Ala Ser 50 60

Ile Met Ser Pro Ile Ala Thr Val Asn Ala Ser Gly Leu Ser Lys Glu Gln Leu Glu His Arg Glu Arg Ser Leu Gln Thr Leu Arg Asp Ile Glu Arg Leu Leu Arg Ser Gly Glu Thr Glu Pro Phe Leu Lys Gly Ala Pro Arg Arg Ser Gly Gly Leu Lys Lys Tyr Glu Glu Pro Leu Gln Ser Met Ile Ser Gln Thr Gln Ser Leu Gly Gly Pro Pro Leu Glu His Glu Val Pro Gly His Pro Pro Gly Gly Asp Met Gly Gln Gln Met Asn Met Met Ile Gln Arg Leu Gly Gln Asp Ser Leu Thr Pro Glu Gln Val Ala Trp Arg Lys Leu Gln Glu Glu Tyr Tyr Glu Glu Lys Arg Arg Lys Glu Glu Gln Ile Gly Leu His Gly Ser Arg Pro Leu Gln Asp Met Met Gly Met Gly Gly Met Met Val Arg Gly Pro Pro Pro Pro Tyr His Ser Lys Pro Gly Asp Gln Trp Pro Pro Gly Met Gly Ala Gln Leu Arg Gly Pro Met Asp Val Gln Asp Pro Met Gln Leu Arg Gly Pro Pro Phe Pro Gly Pro Arg Phe Pro Gly Asn Gln Ile Gln Arg Val Pro Gly Phe Gly Gly Met Gln Ser Met Pro Met Glu Val Pro Met Asn Ala Met Gln Arg

Pro	Val 290	Arg	Pro	Gly	Met	Gly 295	Trp	Thr	Glu	Asp	Leu 300	Pro	Pro	Met	Gly
Gly 305	Pro	Ser	Asn	Phe	Ala 310	Gln	Asn	Thr	Met	Pro 315	Tyr	Pro	Gly	Gly	Gln 320
Gly	Glu	Ala	Glu	Arg 325	Phe	Met	Thr	Pro	Arg 330	Val	Arg	Glu	Glu	Leu 335	Leu
Arg	His	Gln	Leu 340	Leu	Glu	Lys	Arg	Ser 345	Met	Gly	Met	Gln	Arg 350	Pro	Leu
Gly	Met	Ala 355	Gly	Ser	Gly	Met	Gly 360	Gln	Ser	Met	Glu	Met 365	Glu	Arg	Met
Met	Gln 370	Ala	His	Arg	Gln	Met 375	Asp	Pro	Ala	Met	Phe 380	Pro	Gly	Gln	Met
Ala 385	Gly	Gly	Glu	Gly	Leu 390	Ala	Gly	Thr	Pro	Met 395	Gly	Met	Glu	Phe	Gly 400
Gly	Gly	Arg	Gly	Leu 405	Leu	Ser	Pro	Pro	Met 410	Gly	Gln	Ser	Gly	Leu 415	Arg
Glu	Val	Asp	Pro 420	Pro	Met	Gly	Pro	Gly 425	Asn	Leu	Asn	Met	Asn 430	Met	Asn
Val	Asn	Met 435	Asn	Met	Asn	Met	Asn 440	Leu	Asn	Val	Gln	Met 445	Thr	Pro	Gln
Gln	Gln 450	Met	Leu		Ser							Gly	Asp	Leu	Met
Gly 465	Pro	Gln	Gly	Leu	Ser 470	Pro	Glu	Glu	Met	Ala 475	Arg	Val	Arg	Ala	Gln 480
Asn	Ser	Ser	Gly	Met 485	Val	Pro	Leu	Pro	Ser 490	Ala	Asn	Pro	Pro	Gly 495	Pro
Leu	Lys	Ser	Pro 500	Gln	Val	Leu	Gly	Ser 505	Ser	Leu	Ser	Val	Arg 510	Ser	Pro
Thr	Gly	Ser	Pro	Ser	Arg	Leu	Lys	Ser	Pro	Ser	Met	Ala	Val	Pro	Ser

Pro	Gly 530	Trp	Val	Ala	Ser	Pro 535	Lys	Thr	Ala	Met	Pro 540	Ser	Pro	Gly	Val
Ser 545	Gln	Asn	Lys	Gln	Pro 550	Pro	Leu	Asn	Met	Asn 555	Ser	Ser	Thr	Thr	Leu 560
Ser	Asn	Met	Glu	Gln 565	Asp	Pro	Thr	Pro	Ser 570	Gln	Asn	Pro	Leu	Ser 575	Leu
Met	Met	Thr	Gln 580	Met	Ser	Lys	туr	Ala 585	Met	Pro	Ser	Ser	Thr 590	Pro	Leu
Tyr	His	Asn 595	Ala	Ile	Lys	Thr	Ile 600	Ala	Thr	Ser	Asp	Asp 605	Glu	Leu	Leu
Pro	Asp 610	Arg	Pro	Leu	Leu	Pro 615	Pro	Pro	Pro	Pro	Pro 620	Gln	Gly	Ser	Gly
Pro 625	Gly	Gly	Pro	Asp	Ser 630	Leu	Asn	Ala	Pro	Суs 635	Gly	Pro	Val	Pro	Ser 640
Ser	Ser	Gln	Met	Met 645	Pro	Phe	Pro	Pro	Arg 650	Leu	Gln	Gln	Pro	His 655	Gly
Ala	Met	Ala	Pro 660	Thr	Gly	Gly	Gly	Gly 665	Gly	Gly	Pro	Gly	Leu 670	Gln	Gln
His	Tyr	Pro 675	Ser	Gly	Met	Ala	Leu 680	Pro	Pro	Glu	Asp	Leu 685	Pro	Asn	Gln
Pro	Pro 690	Gly	Pro	Met	Pro	Pro 695	Gln	Gln	His	Leu	Met 700	Gly	Lys	Ala	Met
Ala 705	Gly	Arg	Met	Gly	Asp 710	Ala	Tyr	Pro	Pro	Gly 715	Val	Leu	Pro	Gly	Val 720
Ala	Ser	Val	Leu	Asn 725	Asp	Pro	Glu	Leu	Ser 730	Glu	Val	Ile	Arg	Pro 735	Thr
Pro	Thr	Gly	Ile 740	Pro	Glu	Phe	Asp	Leu 745	Ser	Arg	Ile	Ile	Pro 750	Ser	Glu

Lys Pro Ser Ser Thr Leu Gln Tyr Phe Pro Lys Ser Glu Asn Gln Pro Pro Lys Ala Gln Pro Pro Asn Leu His Leu Met Asn Leu Gln Asn Met Met Ala Glu Gln Thr Pro Ser Arg Pro Pro Asn Leu Pro Gly Gln Gln Gly Asp Arg Pro Leu Val Val Val Ile Pro Gly Thr Arg Ala Met Ala Pro Ala Gln Arg Cys Pro Leu Cys Arg Gln Thr Phe Phe Cys Gly Arg Gly His Val Tyr Ser Arg Lys His Gln Arg Gln Leu Lys Glu Ala Leu Glu Arg Leu Leu Pro Gln Val Glu Ala Ala Arg Lys Ala Ile Arg Ala Ala Gln Val Glu Arg Tyr Val Pro Glu His Glu Arg Cys Cys Trp Cys Leu Cys Cys Gly Cys Glu Val Arg Glu His Leu Ser His Gly Asn Leu Thr Val Leu Tyr Gly Gly Leu Leu Glu His Leu Ala Ser Pro Glu His Lys Lys Ala Thr Asn Lys Phe Trp Trp Glu Asn Lys Ala Glu Val Gln Met Lys Glu Lys Phe Leu Val Thr Pro Gln Asp Tyr Ala Arg Phe Lys Lys Ser Met Val Lys Gly Leu Asp Ser Tyr Glu Glu Lys Glu Asp Lys Val Ile Lys Glu Met Ala Ala Gln Ile Arg Glu Val Glu Gln Ser Arg

Gln Glu Val Val Arg Ser Val Leu Glu Thr Gly Pro Pro Arg Tyr Ala 980

Leu Thr Val Arg Ser Pro Ala Val Leu Ser Arg Arg Thr Leu Lys Ser 1000

Gly Ala Phe Pro Pro Gln Thr Pro Glu Ala His Pro Gln Ala Arg 1010 1015

Cys Leu Cys Ala Pro Arg Arg Gly Ala Leu Lys Pro Glu Pro Pro 1030 1035

Gly Arg Thr Leu Lys Leu Gly Val Pro Pro His Thr Thr Arg Lys 1045 1050

Ala Arq Pro His Ala Ala Lys Thr Ser Pro Arg Pro Arg Cys Thr 1060 1055

Arg Gln Ala Pro Asn Lys Thr Gln Ser Leu Gln Leu Ala Gly Lys 1075 1070

Ala Arg Lys Thr Ala Leu His Leu Gln Thr Lys Ala Leu Val Gly 1085 1090

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Asp Leu 1115

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<213> Artificial

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	misc_structure (1)(27) T7 Promoter				
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